

<110> Sheppard, Paul O.
Adler, David A.

<130> 97-71

<151> 1999-03-17

<160> 9

<170> FastSEQ for Windows Version 3.0

 $\langle 210 \rangle$ 1

<211> 1008

<212> DNA

<213> Homo sapiens

 $\langle 220 \rangle$

<221> CDS

<222> (128)...(784)

 $\langle 400 \rangle$ 1

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aatcaaaa atg aag ctt ctc ctt tgg gcc tgc att gta tgt gtt gct ttt      169
      Met Lys Leu Leu Leu Trp Ala Cys Ile Val Cys Val Ala Phe
          1             5             10

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gca agg aag aga cgg ttc ccc ttc att ggt gag gat gac aat gac gat 217
Ala Arg Lys Arg Arg Phe Pro Phe Ile Gly Glu Asp Asp Asn Asp Asp
15 20 25 30

ggt cac cca ctt cat cca tct ctg aat att cct tat ggc ata cgg aat 265
Gly His Pro Leu His Pro Ser Leu Asn Ile Pro Tyr Gly Ile Arg Asn
35 40 45

tta cca cct cct ctt tat tat cgc cca gtg aat aca gtc ccc agt tac 313
Leu Pro Pro Pro Leu Tyr Tyr Arg Pro Val Asn Thr Val Pro Ser Tyr
50 55 60

cct ggg aat act tac act gac aca ggg tta cct tcg tat ccc tgg att	361
Pro Gly Asn Thr Tyr Thr Asp Thr Gly Leu Pro Ser Tyr Pro Trp Ile	
65 70 75	
cta act tct cct gga ttc ccc tat gtc tat cac atc cgt ggt ttt ccc	409
Leu Thr Ser Pro Gly Phe Pro Tyr Val Tyr His Ile Arg Gly Phe Pro	
80 85 90	
tta gct act cag ttg aat gtt cct cct ctc cct cct agg ggt ttc ccg	457
Leu Ala Thr Gln Leu Asn Val Pro Pro Leu Pro Pro Arg Gly Phe Pro	
95 100 105 110	
ttt gtc cct cct tca agg ttt ttt tca gca gct gca gca ccc gct gcc	505
Phe Val Pro Pro Ser Arg Phe Phe Ser Ala Ala Ala Ala Pro Ala Ala	
115 120 125	
cca cct att gca gct gag cct gct gca gct gca cct ctt aca gcc aca	553
Pro Pro Ile Ala Ala Glu Pro Ala Ala Ala Pro Leu Thr Ala Thr	
130 135 140	
cct gta gca gct gag cct gct gca ggg gcc cct gtt gca gct gag cct	601
Pro Val Ala Ala Glu Pro Ala Ala Gly Ala Pro Val Ala Ala Glu Pro	
145 150 155	
gct gca gag gca cct gtt gga gct gag cct gct gca gag gca cct gtt	649
Ala Ala Glu Ala Pro Val Gly Ala Glu Pro Ala Ala Glu Ala Pro Val	
160 165 170	
gca gct gag cct gct gca gag gca cct gtt gga gtg gag cca gct gca	697
Ala Ala Glu Pro Ala Ala Glu Ala Pro Val Gly Val Glu Pro Ala Ala	
175 180 185 190	
gag gaa cct tca cca gct gag cct gct aca gcc aag cct gct gcc cca	745
Glu Glu Pro Ser Pro Ala Glu Pro Ala Thr Ala Lys Pro Ala Ala Pro	
195 200 205	
gaa cct cac cct tct ccc tct ctt gaa cag gca aat cag tgaaattctc	794
Glu Pro His Pro Ser Pro Ser Leu Glu Gln Ala Asn Gln	
210 215	
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ttttctttct tttccaaaga ctatttcatt ctgttgatt cagagtattc atctcactac	914
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<221> misc feature

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